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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/987,899

DATE: 11/21/2001
TIME: 10:09:42

P5

Input Set : F:\sequence listing for 16517.258.txt
Output Set: N:\CRF3\11212001\I987899.raw

2 <110> APPLICANT: Cheikh, Nordine
 3 Liu, Jingdong
 4 Miller, Phillip W.
 5 O Connell, Keith M.
 7 <120> TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
 8 With the Carbon Assimilation Pathway
 10 <130> FILE REFERENCE: 16517.258
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/987,899
 C--> 12 <141> CURRENT FILING DATE: 2001-11-16
 12 <150> PRIOR APPLICATION NUMBER: US 09/262,979
 13 <151> PRIOR FILING DATE: 1999-03-04
 15 <150> PRIOR APPLICATION NUMBER: US 60/076,712
 16 <151> PRIOR FILING DATE: 1998-03-06
 18 <160> NUMBER OF SEQ ID NOS: 7341
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 67
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Zea mays
 25 <220> FEATURE:
 26 <221> NAME/KEY: unsure
 27 <222> LOCATION: (1)...(67)
 28 <223> OTHER INFORMATION: unsure at all n locations
 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Clone ID: 700430856H1
 33 <400> SEQUENCE: 1
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 37 aaaaaaaaaa 67
 40 <210> SEQ ID NO: 2
 41 <211> LENGTH: 142
 42 <212> TYPE: DNA
 43 <213> ORGANISM: Zea mays
 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: Clone ID: 700433144H1
 48 <400> SEQUENCE: 2
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 52 gtgaatgcaa gtcggcata tcatacatat gtaatatggta tacaaactat atatataaaat 120
 54 ccggcgaggc gccgacaata ct 142
 57 <210> SEQ ID NO: 3
 58 <211> LENGTH: 142
 59 <212> TYPE: DNA
 60 <213> ORGANISM: Zea mays
 62 <220> FEATURE:
 63 <223> OTHER INFORMATION: Clone ID: 700433148H1
 65 <400> SEQUENCE: 3
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 69 gtgaatgcaa gtcggcata tcatacatat gtaatatggta tacaaactat atatataaaat 120
 71 ccggcgaggc gccgacaata ct 142

ENTERED

RAW SEQUENCE LISTING
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Input Set : F:\sequence listing for 16517.258.txt
Output Set: N:\CRF3\11212001\I987899.raw

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74 <210> SEQ ID NO: 4
75 <211> LENGTH: 319
76 <212> TYPE: DNA
77 <213> ORGANISM: Zea mays
79 <220> FEATURE:
80 <223> OTHER INFORMATION: Clone ID: 700098783H1
82 <400> SEQUENCE: 4
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86 ggcctcgctg gccaccggcg tcgctccgtt ccaggggctc aagtccaccg ccagccccc 120
88 cgtcgcccgcc cgctcctcca gaagcctcgg caacgtcagc aacggcgaa ggatccggtg 180
90 catgcaggtg tggccggcct acggcaacaa gaagttcag aacgtgtcgat acctgacgcc 240
92 gctgtcgacg gacgacttcg tgaagcaggt ggactacctg ctgcgcacg gctggatacc 300
94 ctgcctcgag ttccatcaag 319
97 <210> SEQ ID NO: 5
98 <211> LENGTH: 323
99 <212> TYPE: DNA
100 <213> ORGANISM: Zea mays
102 <220> FEATURE:
103 <223> OTHER INFORMATION: Clone ID: 700097213H1
105 <400> SEQUENCE: 5
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109 cagcaagggtc ggcttcgtgt accgcgagaa ctccacccctcc ccgtgctact acgacggccg 120
111 ctactggacc atgttggaaatc atgcccattt tcggctgcaa cgacgcacc caggtgtaca 180
113 aggagctgca ggaggccatc aaatccatcc cggacgcctt ccaccgcgtc atccgcttcg 240
115 acaacatcaa gcagacgcag tgcgtcagttt tcacgccta caagccccgg ggcagcgtact 300
117 agaccgcgcgc ccc 323
120 <210> SEQ ID NO: 6
121 <211> LENGTH: 330
122 <212> TYPE: DNA
123 <213> ORGANISM: Zea mays
125 <220> FEATURE:
126 <223> OTHER INFORMATION: Clone ID: 700097673H1
128 <400> SEQUENCE: 6
130 gcttgcctat ggcgcacccg tgatgtatggc ctcgtcgac accgcgcgtc ccccggttcca 60
132 gggctcaag tccgcgcgcgc gcctcccgat ccgcgcgcgc agcaccagga gcctcgccaa 120
134 cgtcagcaac ggcggaaatc tccgggtcat gcaggtgtgg ccggccctacg gcaacaagaa 180
136 gttcgagacg ctgttgttacc tgcgtcgatc gtcgacggac gacctgttgc agcaggttgg 240
138 ctacctgttgc gcaacggctt ggataccctg cctcgatcc agcaagctcg ggttcgttgc 300
140 ccgcgagaac tccacccccc cgtgtacta 330
143 <210> SEQ ID NO: 7
144 <211> LENGTH: 310
145 <212> TYPE: DNA
146 <213> ORGANISM: Zea mays
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Clone ID: 700101767H1
151 <400> SEQUENCE: 7
153 agcaagcaag ctcgtatctac tactactact agctggatca catactagcc agcctgcac 60
155 ccagcttgcgcgcgc atggcgccca ccgtgatgat ggcctcgatc gcaaccggcg tcggcccgat 120
157 ccagggtctc aagtccgcgcgc ccagccccc cgtcgccgc cgcagcacca ggagccctcg 180

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159 caacgtcagc aacggcgaa ggatccggtg catcagggtg tggccggct acggcaacaa 240
161 gaagttcagc acgctgtcgt acctgccccc gctgtcagc gacgacactgc tgaagcagg 300
163 ggactacctg 310
166 <210> SEQ ID NO: 8
167 <211> LENGTH: 318
168 <212> TYPE: DNA
169 <213> ORGANISM: Zea mays
171 <220> FEATURE:
172 <223> OTHER INFORMATION: Clone ID: 700100001H1
174 <400> SEQUENCE: 8
176 gtcgccccgcc gcagcaccag gagcctcggc aacgtcagca acggcgaaag gatccgggtgc 60
178 atgcagggtgt ggccggccta cggcaacaag aagtccgaga cgctgtcgtta cctgcccgg 120
180 ctgtcgacgg acgacctgct gaagcagggtg gactacactgc tgcccaacgg ctggataccc 180
182 tgcctcgagt tcagcaagct cgggttcgtt taccgcgaga actccacactc cccgtgtac 240
184 tacgacggcc gctactggac catgttggaaatgtt ctggctgcac cgacgcccacg 300
186 caagtgtaca aggagct 318
189 <210> SEQ ID NO: 9
190 <211> LENGTH: 316
191 <212> TYPE: DNA
192 <213> ORGANISM: Zea mays
194 <220> FEATURE:
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201 atgatggcct cgtcggccac cggccgtcgtt ccgttccagg ggctcaagtc caccggccagc 120
203 ctccccatcg cccggccgtc ctccagaagc ctcggcaacg tcagcaacgg cggaaggatc 180
205 cgggtcatgc aggtgtggcc ggcttacggc aacaagaagt tcgagacgt gtgttacctg 240
207 cccggctgtt cgacggacga cctgttgaag caggtggact acctgtcgca caacggctgg 300
209 ataccctgcc tcgagt 316
212 <210> SEQ ID NO: 10
213 <211> LENGTH: 317
214 <212> TYPE: DNA
215 <213> ORGANISM: Zea mays
217 <220> FEATURE:
218 <223> OTHER INFORMATION: Clone ID: 700099925H1
220 <400> SEQUENCE: 10
222 agcaagcaag caagcagcgtt gtacatacat actaggcagc caggcagcca tggcgcccac 60
224 cgtatgtatg gcctgtcggtt ccaccggcgtt cgctccgttc caggggctca agtccaccgc 120
226 cagcctccccc gtcggccggcc gtcctccagg aagcctcgcc aacgtcagca acggcgaaag 180
228 gatccgggtgc atgcagggtgt ggccggccata cggcaacaag aagttcgaga cgctgtcgtta 240
230 cctggccggcc ctgtcgacgg acgacctgctt gaagcagggtg gactacactgc tgcccaacgg 300
232 ctggataccc tgcctcg 317
235 <210> SEQ ID NO: 11
236 <211> LENGTH: 315
237 <212> TYPE: DNA
238 <213> ORGANISM: Zea mays
240 <220> FEATURE:
241 <223> OTHER INFORMATION: Clone ID: 700098235H1
243 <400> SEQUENCE: 11

RAW SEQUENCE LISTING
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245	caagcaagca	agcagcgt	acatacatac	taggcagcca	ggcagccatg	gcgcccaccg	60
247	tgtatggc	ctcgccggcc	accgcgcgtc	ctccgttcca	ggggctcaag	tccaccgcca	120
249	gcctccccgt	cgcgcgcgc	tcctccagaa	gcctcgccaa	cgtcagcaac	ggcggaaagga	180
251	tccgggtgc	catgcagggtgg	ccggcctacg	gcaacaagaa	gttcgagacg	ctgtcgtaacc	240
253	tgccgcgcgt	gtcgacggac	gacctgctga	agcaggtgga	ctacctgctg	cgcaacggct	300
255	ggataccctg	cctcg					315
258	<210>	SEQ ID NO:	12				
259	<211>	LENGTH:	302				
260	<212>	TYPE:	DNA				
261	<213>	ORGANISM:	Zea mays				
263	<220>	FEATURE:					
264	<223>	OTHER INFORMATION:	Clone ID: 700093043H1				
266	<400>	SEQUENCE:	12				
268	gcaagctcg	tctactacta	ctactagctg	gtacacatac	tagccagcct	gccagccagc	60
270	ttgccatggc	gcccaccgtg	atgatggcct	cgtcgcaac	cgcgtcgcc	cggttccagg	120
272	gtctcaagtc	cgcgcgcgc	ctccgcgtc	cccgcgcag	caccaggagc	ctcgcaacg	180
274	ttagcaacgg	cggaaaggatc	cggtgcattgc	agggtgtggcc	ggcctacggc	aacaagaagt	240
276	tgcagacgt	gtcgtaacctg	ccggcgttgt	cgacggacga	cctgctgaag	caggtggact	300
278	ac						302
281	<210>	SEQ ID NO:	13				
282	<211>	LENGTH:	315				
283	<212>	TYPE:	DNA				
284	<213>	ORGANISM:	Zea mays				
286	<220>	FEATURE:					
287	<223>	OTHER INFORMATION:	Clone ID: 700089802H1				
289	<400>	SEQUENCE:	13				
291	agcaagcaag	cagcgtgtac	atacatacta	ggcagccagg	cagccatggc	gcccacccgtg	60
293	atgatggcct	cgtcgccac	cgcgcgtcgt	ccgttccagg	ggctcaagtc	cacgcccagcc	120
295	tccgcgtc	ccgcgcgtcc	tccagaagcc	tcggcaacgt	cagcaacggc	ggaaggatcc	180
297	gggtgcattgc	gggtgtggccg	gcctacggc	acaagaagtt	cgagacgtg	tcttacactgc	240
299	cgcgcgttc	gacggacgac	ctgctgaagc	aggtggacta	cctgctgcgc	aacggctgga	300
301	taccctgcct	cgagt					315
304	<210>	SEQ ID NO:	14				
305	<211>	LENGTH:	297				
306	<212>	TYPE:	DNA				
307	<213>	ORGANISM:	Zea mays				
309	<220>	FEATURE:					
310	<223>	OTHER INFORMATION:	Clone ID: 700101196H1				
312	<400>	SEQUENCE:	14				
314	gccaggcagc	catggcgccc	accgtgtatga	tggcctcg	ggccaccggc	gtcgctccgt	60
316	tccaggggct	caagtccacc	gccagcctcc	ccgtcgcccg	ccgtccctcc	agaaggctcg	120
318	gcaacgtcag	caacggcgg	aggatccgt	gcatgcagg	gtggccggcc	tacggcaaca	180
320	agaagttcga	gacgtgtcg	tacctgcgc	cgctgtcgac	ggacgacgt	ctgaaggcagg	240
322	tggactaccc	gtcgcaac	ggctggatac	cctgcctcg	gttcagcaag	gtcggt	297
325	<210>	SEQ ID NO:	15				
326	<211>	LENGTH:	309				
327	<212>	TYPE:	DNA				
328	<213>	ORGANISM:	Zea mays				
330	<220>	FEATURE:					

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Input Set : F:\sequence listing for 16517.258.txt
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331 <223> OTHER INFORMATION: Clone ID: 700097309H1
 333 <400> SEQUENCE: 15
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 337 tggatgtggc ctcgtcgcc accgcgtcg ctccgttcca ggggctcaag tccaccgcca 120
 339 gcctccccgt cgccgcgc tcctccagaa gcctcggcaa cgtcagcaac ggcggaaagga 180
 341 tccggtgcat gcagggtgtgg ccggcctacg gcaacaagaa gttcggagacg ctgtcgtaacc 240
 343 tgccgcgcgt gtcgacggac gacctgtga agcaggtgga ctacctgtcg cacaacggct 300
 345 ggataccct 309
 348 <210> SEQ ID NO: 16
 349 <211> LENGTH: 305
 350 <212> TYPE: DNA
 351 <213> ORGANISM: Zea mays
 353 <220> FEATURE:
 354 <223> OTHER INFORMATION: Clone ID: 700100270H1
 356 <400> SEQUENCE: 16
 358 agcaaggcagc gagtacatac atactaggca gccaggcagc catggcgccc accgtgatga 60
 360 tggcctcgctc ggccacccgccc gtcgctccgt tccaggggtt caagtccacc gccagccccc 120
 362 ccgtcgcccg cccgtctcc agaaggcctcg gcaacgtcag caacggcgga aggatccgg 180
 364 gcatgcagggt gtggccggcc tacggcaaca agaagttcga gacgctgtcg tacctgcgc 240
 366 cgctgtcgac ggacgacctg ctgaaggcagg tggactacct gctgcgcaac ggctggataac 300
 368 cctgc 305
 371 <210> SEQ ID NO: 17
 372 <211> LENGTH: 312
 373 <212> TYPE: DNA
 374 <213> ORGANISM: Zea mays
 376 <220> FEATURE:
 377 <223> OTHER INFORMATION: Clone ID: 700208152H1
 379 <400> SEQUENCE: 17
 381 caacgcgtcc gggacctggg caagcaagct cgatctacta ctactacta ctggtagacaca 60
 383 tactagccag cctgcacccgacc agcttgccat ggcgcaccgt gtgtatgtgg cctcgccggc 120
 385 aaccgcgtcc gcccgttcc agggctccaa gtccgcgc accctcccg tcgcgc 180
 387 cagcaccagg agcctcggca acgtcagccaa cggcgaaagg atccggtgca tgcagggttg 240
 389 gcccgcctac ggcaacaaga agttcgagac gctgtcgatc ctgcgcgcg tgcgacgg 300
 391 cgacactgtaa 312
 394 <210> SEQ ID NO: 18
 395 <211> LENGTH: 322
 396 <212> TYPE: DNA
 397 <213> ORGANISM: Zea mays
 399 <220> FEATURE:
 400 <221> NAME/KEY: unsure
 401 <222> LOCATION: (1)...(322)
 402 <223> OTHER INFORMATION: unsure at all n locations
 404 <220> FEATURE:
 405 <223> OTHER INFORMATION: Clone ID: 700215709H1
 407 <400> SEQUENCE: 18
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 411 agccagctt ccatggccaccgtatggctcgatg atggcctcgatg cggcaaccgc cgtcgccccc 120
 413 ttccagggttc tcaagtccgc cggcgcctc cccgtcgccc gccgcagcac caggagcctc 180
 415 ggcaacgtca gcaacggcg aaggatccgg tgcatgcagg tgtggccggc ctacggcaac 240

Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/987,899

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Input Set : F:\sequence listing for 16517.258.txt
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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:35 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68
L:1498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68
L:2159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100
L:2204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102
L:2206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102
L:2208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102
L:2329 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:108
L:2391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:111
L:2395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:111
L:2609 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2611 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2613 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2615 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2617 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2619 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2621 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122
L:2704 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:126
L:2747 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:128
L:2831 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2833 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2835 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2837 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2839 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2841 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2843 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2845 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:2863 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132
L:2865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132
L:2867 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132
L:2873 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132
L:2875 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132
L:2893 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:133
L:2969 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:136
L:3016 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3018 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3020 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3022 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3024 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3026 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3028 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138
L:3048 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139
L:3050 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139
L:3052 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139

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L:3054 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139
L:3056 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139
L:3058 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139
L:3060 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139